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40 45 Thr Thr Gly Lys Leu Pro Val Pro Trp Pro Thr Leu Val Thr Thr Leu Ser Tyr Gly Val Gln Cys Phe Ser Arg Tyr Pro Asp His Met Lys Gln His Asp Phe Phe Lys Ser Ala Met Pro Glu Gly Tyr Ile Gln Glu Arg 90 Thr Ile Phe Phe Glu Asp Asp Gly Asn Tyr Lys Ser Arg Ala Glu Val 105 Lys Phe Glu Gly Asp Thr Leu Val Asn Arg Ile Glu Leu Thr Gly Thr 120 Asp Phe Lys Glu Asp Gly Asn Ile Leu Gly Asn Lys Met Glu Tyr Asn 135 Tyr Asn Ala Ser Asn Val Tyr Ile Met Thr Asp Lys Ala Lys Asn Gly Ile Lys Val Asn Leu Lys Ile Arg His Asn Ile Ala Asp Gly Ser Val 170 175 165 Gln Leu Ala Asp His Tyr Gln Gln Asn Thr Pro Ile Gly Asp Gly Pro 185 Val Leu Leu Pro Asp Asn His Tyr Leu Ser Thr Gln Ser Ala Leu Ser 200 Lys Asp Pro Asn Glu Lys Arg Asp His Met Ile Tyr Phe Glu Phe Val 215 220 Thr Ala Ala Ile Thr His Gly Met Asp Glu Leu Ile Lys <210> 23 <211> 717 <212> DNA <213> Aequoria coerulescens <400> 23 atgageaagg gegeegaget gtteaeegge ategtgeeca teetgatega getgaatgge 60 qatqtqaatq gccacaaqtt cagcgtgagc ggcgagggcg agggcgatgc cacctacggc 120 aagetgacce tgaagtteat etgeaceace ggeaagetge etgtgeeetg geeeaceetg 180 gtgaccaccc tgagctacgg cgtgcagtgc ttctcacgct accccgatca catgaagcag 240 cacgacttct tcaagagcgc catgcctgag ggctacatcc aggagcgcac catcttcttc 300 gaggatgacg gcaactacaa gtcgcgcgcc gaggtgaagt tcgagggcga taccctggtg 360 aatcgcatcg agctgaccgg caccgatttc aaggaggatg gcaacatcct gggcaataag 420 atggagtaca actacaacgc ccacaatgtg tacatcatga ccgacaaggc caagaatggc 480 atcaaggtga acttcaagat ccgccacaac atcgaggatg gcagcgtgca gctggccgac 540 cactaccage agaatacccc categgegat ggccetgtge tgetgeeega taaccactae 600 ctgtccaccc agagcgccct gtccaaggac cccaacgaga agcgcgatca catgatctac 660 tteggetteg tgaeegeege egecateace caeggeatgg atgagetgta caagtga <210> 24 <211> 238 <212> PRT <213> Aequoria coerulescens <400> 24 Met Ser Lys Gly Ala Glu Leu Phe Thr Gly Ile Val Pro Ile Leu Ile Glu Leu Asn Gly Asp Val Asn Gly His Lys Phe Ser Val Ser Gly Glu

Gly Glu Gly Asp Ala Thr Tyr Gly Lys Leu Thr Leu Lys Phe Ile Cys

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<211> 238

<212> PRT

<213> Aequoria coerulescens

<400> 25

Met Ser Lys Gly Glu Glu Leu Phe Thr Gly Val Val Pro Ile Leu Val Glu Leu Asp Gly Asp Val Asn Gly His Lys Phe Ser Val Ser Gly Glu 25 Gly Glu Gly Asp Ala Thr Tyr Gly Lys Leu Thr Leu Lys Phe Ile Cys Thr Thr Gly Lys Leu Pro Val Pro Trp Pro Thr Leu Val Thr Thr Phe Ser Tyr Gly Val Gln Cys Phe Ser Arg Tyr Pro Asp His Met Lys Gln His Asp Phe Phe Lys Ser Ala Met Pro Glu Gly Tyr Val Gln Glu Arg 85 90 Thr Ile Phe Phe Lys Asp Asp Gly Asn Tyr Lys Thr Arg Ala Glu Val 105 Lys Phe Glu Gly Asp Thr Leu Val Asn Arg Ile Glu Leu Lys Gly Ile 120 Asp Phe Lys Glu Asp Gly Asn Ile Leu Gly His Lys Leu Glu Tyr Asn 135 140 Tyr Asn Ser His Asn Val Tyr Ile Met Ala Asp Lys Gln Lys Asn Gly 150 155 Ile Lys Val Asn Phe Lys Ile Arg His Asn Ile Glu Asp Gly Ser Val 170 Gln Leu Ala Asp His Tyr Gln Gln Asn Thr Pro Ile Gly Asp Gly Pro 180 185

Val	Leu	Leu	Pro	Asp	Asn	His	Tyr	Leu	Ser	Thr	Gln	Ser	Ala	Leu	Ser	
		195					200					205				
Lys	Asp	Pro	Asn	Glu	Lys	Arg	Asp	His	Met	Val	Leu	Leu	Glu	Phe	Val	
	210					215					220					
Thr	Ala	Ala	Gly	Ile	Thr	His	Gly	Met	Asp	Glu	Leu	Tyr	Lys			
225					230					235						